

SEQUENCE LISTING

<110> Kapoor, Archana Munshi, Anil

<120> MEMBRANE-ASSOCIATED IMMUNOGENS OF MYCOBACTERIA

<130> A-57004-4/RFT/JJD

<140> US 09/432,820

<141> 1999-11-02

<150> US 09/099,902

<151> 1998-06-18

<150> US 08/710,676

<151> 1996-09-23

<150> US 08/192,632

<151> 1994-02-07

<150> US 07/906,395

<151> 1992-06-29

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aat cac teg eec ate gag ege age gge ata tgg eag geg etg etg gga 1110 Asn His Ser Pro Ile Glu Arg Ser Gly Ile Trp Gln Ala Leu Leu Gly age gat get att tat tte gag gte geg ggg ggt gte aeg gtg tte gtg 1158 Ser Asp Ala Ile Tyr Phe Glu Val Ala Ala Gly Val Thr Val Phe Val 205 210 215 ctg gtg ggg cgg tat ttc gag gcg cgc gcc aag tcg cag gcg ggc agt 1206 Leu Val Gly Arg Tyr Phe Glu Ala Arg Ala Lys Ser Gln Ala Gly Ser 220 225 230 geg etg aga gee ttg geg geg etg age gee aag gaa gta gee gte etg 1254 Ala Leu Arg Ala Leu Ala Ala Leu Ser Ala Lys Glu Val Ala Val Leu 235 240 cta ccg gat ggg tcg gag atg gtc atc ccg gcc gac gaa ctc aaa gaa 1302 Leu Pro Asp Gly Ser Glu Met Val Ile Pro Ala Asp Glu Leu Lys Glu 250 255 cag cag cgc ttc gtg gtg cgt cca ggg cag ata gtt gcc gcc gac ggc 1350 Gln Gln Arg Phe Val Val Arg Pro Gly Gln Ile Val Ala Ala Asp Gly etc gec gtc gac ggg tec get geg gtc gac atg age geg atg ace gge 1398 Leu Ala Val Asp Gly Ser Ala Ala Val Asp Met Ser Ala Met Thr Gly gag gcc aaa ccg acc cgg gtg cgt ccg ggg ggg cag gtc atc ggc ggc 1446 Glu Ala Lys Pro Thr Arg Val Arg Pro Gly Gly Gln Val Ile Gly Gly 305 acc aca gtg ctt gac ggc cgg ctg atc gtg gag gcg gcc gcg gtg ggc 1494 Thr Thr Val Leu Asp Gly Arg Leu Ile Val Glu Ala Ala Ala Val Gly 320 gee gae ace cag tte gee gga atg gte ege ete gtt gag caa geg cag 1542 Ala Asp Thr Gln Phe Ala Gly Met Val Arg Leu Val Glu Gln Ala Gln 330 335 345 gcg caa aag gcc gac gca cag cga cta gcc gac cgg atc tcc tcg gtg 1590 Ala Gln Lys Ala Asp Ala Gln Arg Leu Ala Asp Arg Ile Ser Ser Val 350 360 ttt gtt ccc gct gtg ttg gtt atc gcg gca cta acc gca gcc gga tgg 1638 Phe Val Pro Ala Val Leu Val Ile Ala Ala Leu Thr Ala Ala Gly Trp 365 cta atc gcc ggg gga caa ccc gac cgt gcc gtc tcg gcc gca ctc gcc 1686 Leu Ile Ala Gly Gly Gln Pro Asp Arg Ala Val Ser Ala Ala Leu Ala 385

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11

											atc Ile					2358
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_			_	_	_	_			_	_	gta Val					2550
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_						_				_	gcc Ala	_	-		_	2646
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	_			_	_		_	_			gly aaa	_		_		2742
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Gly Thr Arg Val Ala Thr Ile Asp Thr Ser Glu Ala Val Asp Ala Ala 50 55 60

Ala Leu Cys Gln Ala Val Arg Arg Ala Gly Tyr Gln Ala Asp Leu Cys 65 70 75 80

Thr Asp Asp Gly Arg Ser Ala Ser Asp Pro Asp Ala Asp His Ala Arg 85 90 95

Gln Leu Leu Ile Arg Leu Ala Ile Ala Ala Val Leu Phe Val Pro Val
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Ala Asp Leu Ser Val Met Phe Gly Val Val Pro Ala Thr Arg Phe Thr 115 120 125

Gly Trp Gln Trp Val Leu Ser Ala Leu Ala Leu Pro Val Val Thr Trp 130 135 140

Ala Ala Trp Pro Phe His Arg Val Ala Met Arg Asn Ala Arg His His 145 150 155 160

Ala Ala Ser Met Glu Thr Leu Ile Ser Val Gly Ile Thr Ala Ala Thr 165 170 175

Ile Trp Ser Leu Tyr Thr Val Phe Gly Asn His Ser Pro Ile Glu Arg 180 185 190

Ser Gly Ile Trp Gln Ala Leu Leu Gly Ser Asp Ala Ile Tyr Phe Glu 195 200 205

Val Ala Ala Gly Val Thr Val Phe Val Leu Val Gly Arg Tyr Phe Glu 210 215 220

Ala Arg Ala Lys Ser Gln Ala Gly Ser Ala Leu Arg Ala Leu Ala Ala 225 230 235 240

Leu Ser Ala Lys Glu Val Ala Val Leu Leu Pro Asp Gly Ser Glu Met 245 250 255

Val Ile Pro Ala Asp Glu Leu Lys Glu Gln Gln Arg Phe Val Val Arg
260 265 270

Pro Gly Gln Ile Val Ala Ala Asp Gly Leu Ala Val Asp Gly Ser Ala 275 280 285

Ala Val Asp Met Ser Ala Met Thr Gly Glu Ala Lys Pro Thr Arg Val 290 295 300

Arg Pro Gly Gly Gln Val Ile Gly Gly Thr Thr Val Leu Asp Gly Arg 305 310 315 320

Leu Ile Val Glu Ala Ala Ala Val Gly Ala Asp Thr Gln Phe Ala Gly 325 330 335

Met Val Arg Leu Val Glu Gln Ala Gln Ala Gln Lys Ala Asp Ala Gln . 340 345 350

Arg Leu Ala Asp Arg Ile Ser Ser Val Phe Val Pro Ala Val Leu Val 355 360 365

Ile Ala Ala Leu Thr Ala Ala Gly Trp Leu Ile Ala Gly Gly Gln Pro 370 375 380

Asp Arg Ala Val Ser Ala Ala Leu Ala Val Leu Val Ile Ala Cys Pro 385 390 395 400

Cys Ala Leu Gly Leu Ala Thr Pro Thr Ala Met Met Val Ala Ser Gly
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Arg Gly Ala Gln Leu Gly Ile Phe Leu Lys Gly Tyr Lys Ser Leu Glu
420 425 430

Ala Thr Arg Ala Val Asp Thr Val Val Phe Asp Lys Thr Gly Thr Leu 435 440 445

Thr Thr Gly Arg Leu Gln Val Ser Ala Val Thr Ala Ala Pro Gly Trp 450 455 460

Glu Ala Asp Gln Val Leu Ala Leu Ala Ala Thr Val Glu Ala Ala Ser 465 470 475 480

Glu His Ser Val Ala Leu Ala Ile Ala Ala Ala Thr Thr Arg Arg Asp 485 490 495

Ala Val Thr Asp Phe Arg Ala Ile Pro Gly Arg Gly Val Ser Gly Thr
500 505 510

Val Ser Gly Arg Ala Val Arg Val Gly Lys Pro Ser Trp Ile Gly Ser 515 520 525

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Leu 545	Gly	Glu	Thr	Ala	Val 550	Phe	Val	Glu	Val	Asp 555	Gly	Glu	Pro	Cys	Gl ₃ 560
Val	Ile	Ala	Val	Ala 565	Asp	Ala	Val	Lys	Asp 570	Ser	Ala	Arg	Asp	Ala 575	
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His	Leu	Asp 675	Val	Val	Pro	Leu	Ala 680	Leu	Asp	Leu	Ala	Arg 685	Ala	Thr	Met
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